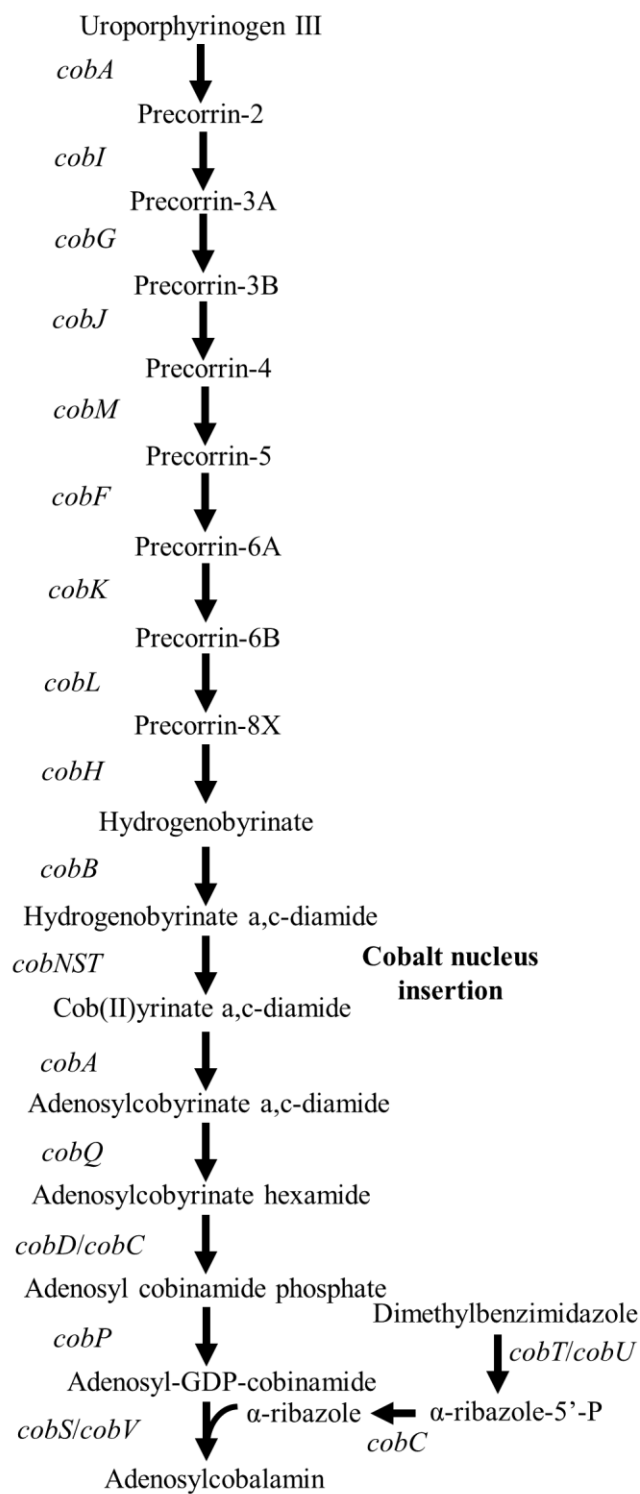


Supplementary Figure S1. 16S rDNA phylogenetic tree for reference (black) and selected (blue) bacteria inferred from Maximum Likelihood (100 bootstrap). Bootstrap values superior to 0.5 are indicated.



Supplementary Figure S2. Vitamin B₁₂ biosynthetic pathway in the sequenced bacteria, starting from uroporphyrinogen III to adenosylcobalamin.